

Amendments to the Specification:

Please amend Table 3 on page 33 as follows:

Table 3. Inferred Amino Acid Sequence of the R2 Envelope Clone from Donor 2

Amino Acid Residue ²					Residue
MRVKGIRRN	QHWWGWGTML	LGLLMICSAT	EKLWVTVYYG	VPVWKEATT	50
LFCASDAKAY	DTEAHNVWAT	HACVPTDPNP	QEVELVNTE	NFNMWKNNMV	100
EQMHEDIISL	WDQSLKPCVK	LTPLCVTLNC	TDLRNTTNTN	NSTDNNNSNS	150
EGTIKGGE MK	N CSFNAITSI	GDKMQKEYAL	LYKLDIEPID	NDNTSYRLIS	200
CNTSVITQAC	PKISFEPIDI	HYCAPAGFAI	LKCNDKKFSG	KGSCKNVSTV	250
QCTHGIRPVV	STQLLLNGSL	AAAAEVVIRSE	N FTNNNAKTI	VQLREPVKIN	300
CSRPNNNTRK	SIPMGPGRAF	YTTGQIIGDI	RQAHCNISK	NWTNALKQVV	350
EKLGEQFNKT	KIVFTNSSGG	DPEIVTHSFN	CAGEFFYCNT	TQLFDIWN	400
ENGTWNITRG	LNNTGRNDT	TLPCRIKQII	NRWQEVGKAM	YAPPIGNIS	450
CSSNITGLL	TRDGGKDDNS	RDGNETFRPG	GGDMRDNWRS	ELYKYKVVKI	500
EPLGVAPTKA	KRRVVQREER	AVGLGAMFIG	FLGAAGSTMG	AASVTLTVQA	550
RQLLSGIVQQ	QSNLLRAIEA	QQHLLQLTVW	GIKQLQARIL	AVERYLKDQQ	600
LLGIWGCSKG	LICTTVPWN	ASWSKNKTLE	AIWNNMTWMQ	WDKEIDNYTK	650
				<u>WDKEIDNYTS</u>	
LIYSLIEESQ	IQQEKNEQEL	LELDKWANKW	NWFDISNWLW	YIKIFIMIVG	700
GLVGLRIVFV	VLSIVNRVRQ	GYSPLSFQTR	LPAPRGPD	EEIEEEGGDR	750
DRDRSGLLVD	GFLTLIWVDL	RSLCLFSYHR	LRDLLLIVTR	IVELLGRRGW	800
EILKYWWNLL	QYWSQELKNS	AVSLFNATAI	AVAEGTDRVI	EVLQRVGRAL	850
LHIPTRIRQG	LERALL				866

²Amino Acid Residues are identified by standard single letter designations. Predicted N-linked glycosylation sites are indicated by shading and bolding